

## THIR UNITED STATES OF AMERICA

The Regents of the Unibersity of California

MICCOM, THERE HAS BEEN PRESENTED TO THE

#### Secretary of Agriculture

AN APPLICATION REQUESTING A CERTIFICATE OF PROTECTION FOR AN ALLEGED DISTINCT VARIETY OF SEXUALLY REPRODUCED, OR TUBER PROPAGATED PLANT. THE NAME AND DESCRIPTION OF WHICH ARE CONTAINED IN THE APPLICATION AND EXHIBITS, A COPY OF WHICH IS HEREUNTO ANNEXED AND MADE A PART HEREOF, AND THE VARIOUS REQUIREMENTS OF LAW IN SUCH GASES MADE AND PROVIDED HAVE BEEN COMPLIED WITH, AND THE TITLE THERETO IS FROM THE RECORDS OF THE PLANT VARIETY PROTECTION OFFICE, IN THE APPLICANT(S) INDICATED IN THE SAID COPY, AND WHEREAS, UPON DUE EXAMINATION MADE, THE SAID APPLICANT(S) IS (ARE) ADJUDGED TO BE ENTITLED TO A CERTIFICATE OF PLANT VARIETY PROTECTION UNDER THE LAW.

NOW, THEREFORE, THIS CERTIFICATE OF PLANT VARIETY PROTECTION IS TO GRANT UNTO THE SAID APPLICANT(S) AND THE SUCCESSORS, HEIRS OR ASSIGNS OF THE SAID APPLICANT(S) FOR THE TERM OF TWENTY HARS FROM THE DATE OF THIS GRANT, SUBJECT TO THE PAYMENT OF THE REQUIRED FEES AND PERIODIC UPLENISHMENT OF VIABLE BASIC SEED OF THE VARIETY IN A PUBLIC REPOSITORY AS PROVIDED BY LAW, THE GHT TO EXCLUDE OTHERS FROM SELLING THE VARIETY, OR OFFERING IT FOR SALE, OR REPRODUCING IT, OR ORITING IT, OR EXPORTING IT, OR CONDITIONING IT FOR PROPAGATION, OR STOCKING IT FOR ANY OF THE PURPOSES, OR USING IT IN PRODUCING A HYBRID OR DIFFERENT VARIETY THEREFROM, TO THE EXTENT OR BY THE PLANT VARIETY PROTECTION ACT. IN THE UNITED STATES SEED OF THIS VARIETY AS A CLASS OF CERTIFIED SEED AND (2) SHALL CONFORM TO THE NUMBER OF THE RIGHTS. (84 STAT. 1542, AS AMENDED, 7 U.S.C. 2321 ET SEO.

BEAN, FIELD

'Canario 707'

In Testimonn Murrest, I have hereunto set my hand and caused the seal of the Plant Haristy Protection Office to be affixed at the City of Washington, D.C. this sixteenth day of Way, in the year two thousand and eight.

Colmonal + Johnfo

Attest:

Bergn

Commissioner Plant Variety Protection Office Secreta Lulture

#### INSTRUCTIONS

GENERAL:Tobeeffectively filedw iththePI antVariety ProtectionO ffice(PVPO), ALLofthefollowi ngitemsmustbereceiv ed inthePVPO:(1)Completed applicationformsignedby theowner;(2)completedexhibitsA,B,C,E;(3)foraseedreproducedvariety atleast2,500viableuntreatedseeds,forahy varietyatleast2,500untreatedseedsofeachlinenecessary toreproduce the variety, or fortuber reproduced varieties verification that a via ble (in the sense that itwillreproduceanentireplant)tissueculturew illbedepositedandmaintainedinanapprovedpublic repository;(4)checkdraw nonaU.S.bankfor\$3,652(\$432 filingfeeand\$3,220examinationfee),pay able to "Treasurer of the United States" (See Section 97.6 of the Regulations and Rules of Practice.) Partial above the Section 100 of the Regulation 10applicationswillbeheldinthePVPO fornotmorethan90day s, thenreturnedtotheapplicantasunfiled.Mailapplicationa ndotherrequirementstoPlantVariety ProtectionOffice, AMS, USDA, Room401, NALBuilding, 10301 Baltimore Avenue, Beltsville, MD 20705-2351.Retainonecopy fory ourfiles.Allitemsonthe faceoftheapplicationareselfexplanatory unlessnotedbelow. Correctionsontheapplicationformandexhibitsmustbeinitialedanddated. **DONOT**use maskingmaterialstomakecorrections. Ifacertificate is a llowed y ouw illberequestedtosendacheckpay ableto"TreasureroftheUnitedStates"intheamount of\$432forissuanceofthecertificate.Certif icatesw illbeissuedtoowner,notlicenseeoragent.

#### PlantVariety ProtectionOffice Telephone:(301)504-5518 FAX:(301)504-5291

Homepage: http://www.ams.usda.gov/science/pvpo/pvp.htm

ITEM

18a. Give:

(1)thegenealogy including publicand commercial varieties, lines, or clones used, and the breeding method;

.(2)thedetailsofsubsequentstagesofselectionandmultiplication;

(3)evidenceofuniformity and stability; and

(4)thety peandfrequency of variants during reproduction and multiplication and state how these variants may

beidentified

18b. Giveasummary ofthevariety 'sdistinctness.Clearly statehow thisapplicationvariety maybedistinguishedfromallothervarietiesinthesamecrop.lfthe newvariety ismostsimilartoonevariety oragroupofrelatedvarieties:

(1)identify thesevarietiesandstatealldifferencesobjectively

(2)attachstatisticaldataforcharactersexpressednumerically

(3)submit,ifhelpful,seedandplantspecimensorphotographs(p

anddemonstratethatthesearecleardifferences; and

rints)ofseedandplantcompar isonsw hichclearly indicatedistinctness.

18c. ExhibitCformsareavailablefromthePVPOOfficeformostcrops;specify aspossibletodescribey ourvariety.

cropkind.FillinEx

hibitC(O bjectiveDescriptionofVariety )formascompletely

18d. Optionaladditionalcharacte risticsand/orphotographs.Describeany additional characteristicsthatcannotbeaccurately convey edinExhibitC.Use comparativevarietiesasisnecessary torevealmoreaccurately thecharacteristicsthatarediffi culttodescribe, such asplanthabit, plantcolor, disease resistance, etc.

18e. Section52(5)oftheActrequiresapplicantstofurnishastatementofthebasisoftheapplicant'sownership.AnExhibitEformisavailablefromthePVPO.

19. If"Y es"isspecified (seedofthisvarietybesoldbyvarietynam eonly, asaclassofcertifiedseed), the applicant MA YNOTreversethisaffirmativedecision afterthevariety hasbeensoldandsolabeled, the decision published, or the certificate issued. How ever,if"No"hasbeenspecified,theapplicantmay changethechoice. (SeeRegulationsandRulesofPractice,Section97.103).

SeeSections41,42, and 43 of the Act and Section 97.5 of the regulations for eligibility requirements.

23. SeeSection55oftheActforinstructionsonclaimi

ngthebenefitofanearlierfilingdate.

21.CONTINUED FROMFRONT

(Pleaseprovideastatementastothelimitati

onandsequenceofgenerationsthatmaybecertified.)

22.CONTINUED FROMFRONT

(Pleaseprovidethedateoffirstsale,

disposition,transfer,oruseforeachcountryandthecircum

stances, if the variety

(includinganyharvestedmaterial)

orahybridproducedfrom thisvarietyhasbeen

sold, disposed of, transferred, or used int

heU.S.orothercountries.)

23.CONTINUEDFROMFRONT

(Pleasegivethecountry, date of filing or issuance, and assigned reference num right(PlantBreeder'sRightorPatent).)

ber, ifthevarietyoranycomponentofthe

varietyisprotectedbyintellectualproperty

NOTES: Itistheresponsibility oftheapplicant/ownertokeepthePVPOinformedofany changesofaddressorchangeofownershiporassignmentorow

ner's

representativeduringthelifeoftheapplication/certificate. assignmentorary modificationofowner'snameisspecifiedinSection97.175oftheregulations.

Thereisnochargeforfilingachangeofaddress. Thefeefor

filingachangeofow (SeeSection101oftheAct, and Sections 97.130, 97.131,

97.175(h)oftheRegulationsandRulesofPractice.)

Toav oidconflictw ithotherv arietynamesinuse,theapplicantmustchecktheappropriaterecognizedauthority .Forexample, foragriculturalanvegetablecrops,contact:SeedBranch,A MS,USDA ,Room213, Building306,Beltsv illeA griculturalResearchCenter—East,Beltsv ille,MD20705. .Forexample, foragriculturaland Telephone:(301)504-8089.http://www.ams.usda.gov /lsg/seed.htm

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Tofi leacom plaintofdiscrimination,writeUSDA,D irector,OfficeofCivilRig hts,R oom326-W ,W hittenBuil ding,14thandindependenceAvenue,SW ,W ashington,DC20250-9410orcali202-720-5964(voiceand TDD).U SDAisanagual opportunitypr oviderandem ployer.

ST-470(02-10-2003)designedby thePlantVar ietyProtectionO fficewithW ord2000.Replac esfor merver sionsofST-470,while h sreobs olete.

## Exhibit A. (Revised November 8, 2007) Origin and Breeding History

UC Canario 707 is a selected and purified, true-breeding variety of *Phaseolus vulgaris* dry bean. The original cross generating "CAP 7" (an early-generation source of Canario 707) was performed in 1984, in the Bean Breeding program of Dr. Steve Temple at the Centro Internacional de Agricultura Tropical (CIAT), Palmira, Colombia. The pedigree of the cross was "84 VA-909 x PAD 2." Parental line "84 VA-909" is a CIAT early-generation breeding line and "PAD 2" is an advanced line derived from the cross G6616 x (G4523 X (G4523 X G76)). Names and origins of all "G" lines are available from the CIAT publication "Catalogo de Germoplasma de Frijol Comun" (October, 1992).

In 1992, the F6 family CAP 7 was planted in the University of California Davis (UC Davis) "Observation" nursery at Chico, California. Like most tropically bred CIAT families, CAP 7 proved to be variable for flowering date, podset, grain size and grain shape, grain color, and maturity. Five plants with the best combination of pod set, erect plant architecture, absence of symptoms of Bean Common Mosaic Virus (BCMV), and large yellow grain, were selected and the seeds were bulked. The bulked seeds were used in a 1993 planting, and from 1993-1996 the line was successively mass selected and bulk harvested at Davis, the Kearney Ag Center (Parlier), West Side Research and Extension Center (Five Points), and Chico. The CAP 7 line was repeatedly mass selected for maturity, plant type, grain type, and pod set. During 1993-1996 testing and mass selection, the selected CAP 7 occasionally showed a low frequency of plants infected with BCMV under field conditions.

Following several seasons of mass selection and bulk harvest, thirty plants were selected and individually harvested from the 1996 UC Davis nursery. The necrosis-inducing NL-3 strain of BCMNV was used to test for the presence of the dominant I gene (F. Morales. *Present status of controlling Bean common mosaic virus, in Plant Virus Disease Control* 524 (A. Hadidi, R. K. Khetarpal and H. Koganezawa, eds., 1998)). Twenty-eight families displayed uniform hypersensitive resistance, and two families segregated for the recessive genotype, ii (BCMV-susceptible). The two families segregating susceptible plants were eliminated, and the resistant twenty-eight families were used to produce UC Canario 707. From each family, 8-10 of the best plants were bulk harvested to produce Breeder seed of UC Canario 707.

The resultant selection process has produced a stable and uniform variety. UC Canario 707 was observed to be uniform and stable over six generations. That is, Breeder and Foundation seed of UC Canario 707 has been observed over six generations and shows stable, uniform expression of grain size, grain color, and hypersensitive resistance to Bean Common Mosaic Virus (BCMV). In June 2002, a random sample of 240 seeds from Breeder seed stock were inoculated with the NL-3 strain, to recheck the stability and uniformity of BCMV resistance. One hundred percent of Canario 707 seedlings tested positive for the dominant I resistance gene. No variants have been observed.

#### Exhibit B (Revised December 27, 2007)

#### **Statement of Distinctness**

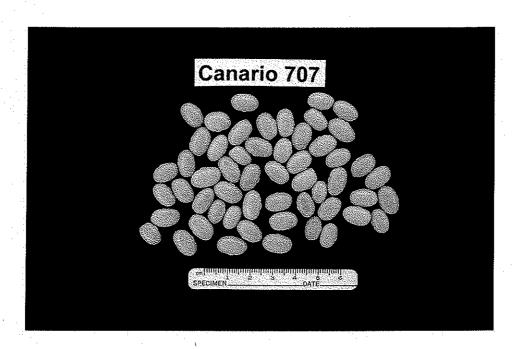
The principal distinguishing features of UC Canario 707 are the combination of a large-seeded, sulfur-yellow grain with I-gene hypersensitive resistance to all identified strains of Bean Common Mosaic Virus (BCMV). Canario 707 grain, examined 2-5 months after harvest, most closely matches hue 7.5Y from the Munsell matte collection (MUNSELL BOOK OF COLOR). Within hue 7.5Y, individual seeds most frequently match chips 7.5Y 8/4, 7.5Y 8/6, 7.5Y 8.5/2, 7.5Y 8.5/4, and 7.5Y 8.5/6.

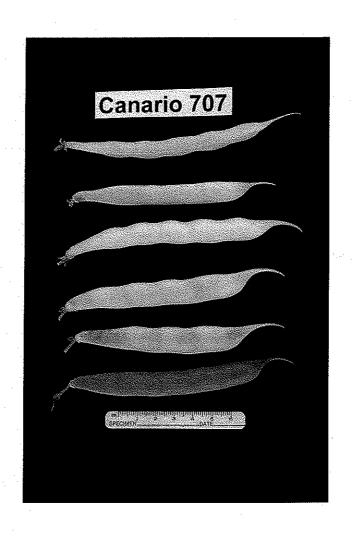
In plant growth habit and grain type, Canario 707 most closely resembles the Mexican cultivar "Mayocoba" (CIAT germplasm accession G 13094) and several yellow-seeded Mexican introductions referred to generically as "Peruanos" (herein referred to as Azufrado Peruano 87). While Canario 707 has resistance to BCMV, conferred by the I gene, Peruano and Mayocoba carry the i allele, conferring susceptibility to some or all strains of BCMV. Further distinguished from Peruano and Mayocoba, Canario 707 displays a hypersensitive response to necrosis-inducing strains of the closely related Bean Common Mosaic Necrosis Virus (BCMNV). F. Morales. Present status of controlling Bean common mosaic virus, in Plant Virus Disease Control 524 (A. Hadidi, R. K. Khetarpal and H. Koganezawa, eds., 1998). Over the six subsequent generations that Canario 707 has been multiplied, resistance to BCMV is stable, uniform, and heritable.

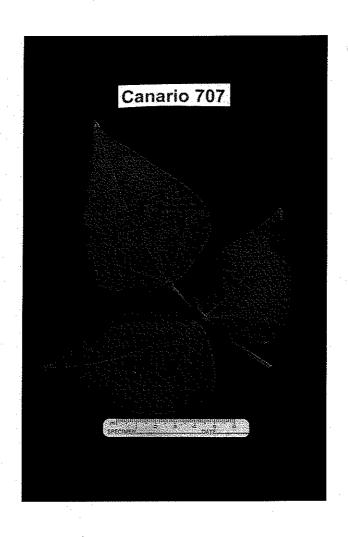
In a comparison with Mayocoba G 13094, Canario 707 seeds have a larger size, greater uniformity, and deeper yellow color. See Figure 1.

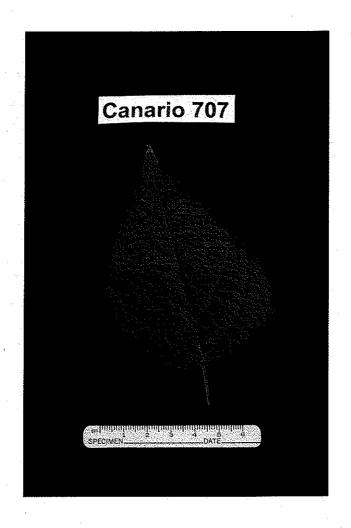
While Canario 707 resembles yellow-seeded varieties, Mayocoba, Peruanos, and Canario 707 varieties can be differentiated by DNA fingerprinting. As indicated in as-filed Exhibit D, Amplified Fragment Length Polymorphism (AFLP) markers were used to fingerprint yellow bean cultivars for determining the genetic relationships between yellow bean and other bean germplasm. AFLP analysis was performed as described by P. Vos et al., Nucleic Acids Research 23: 4407 (1995), and modified by Barcaccia et al., Plant Breed 118: 335 (1999). The primer combinations included five EcoRI-MseI (with selective bases CAC/AAG, CAC/AGC, CCA/AGA, CCA/AGC, and CAA/AAG) and five PstI-MseI combinations (AG/CAC, AG/CAT, AG/CCA, AT/CAA, and AT/CAC). The 10 AFLP primer combinations revealed 133 polymorphic amplified fragments in sampled Andean and Mesoamerican accessions.

Mayocoba (G13094) Canario 707









4-1

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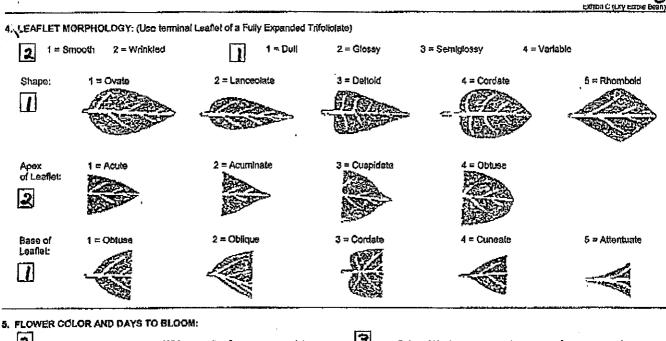
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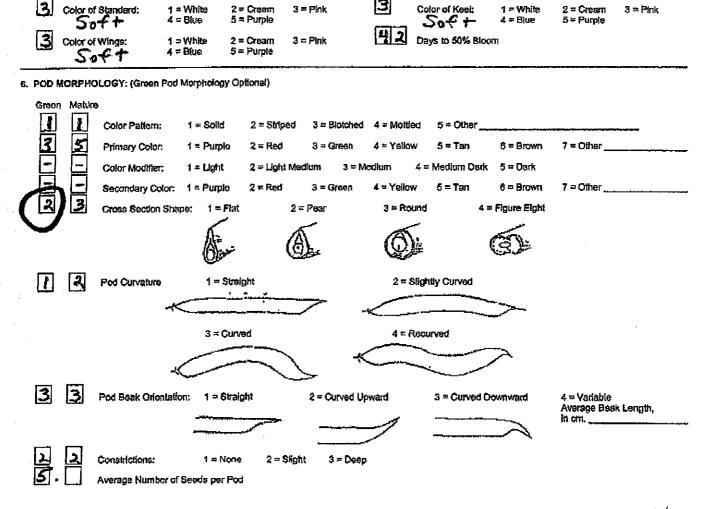
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U.S. DEPARTMENT OF AGRICULTURE AGRICULTURAL MARKETING SERVICE SCIENCE AND TECHNOLOGY PLANT VARIETY PROTECTION OFFICE BELTSVILLE, MD 20705 Exhibit C

## OBJECTIVE DESCRIPTION OF VARIETY Field Bean (Phaseolus vulgaris L.)

•	14/4 47/1 (1 1100000000000000000000000000000000	94.12			
Stove Temple	TEMPORARY OR EXTEREMENTAL RESIGNATIO  Canario 707	Dry Bean			
Plant Sciences De Univ. California Davis, CA, 9561	ot, Mail Stop On	e Prionumber Aprlication No. 200300257			
PLEASE READ ALL INSTRUCTIONS CAREFULLY:					
Provide data for all characters unless indicated as "optio Measured data should be the mean of an appropriate nu may be used to determine plant color. Designate the col	mber of well spaced (15-20 cm) blants.	characters or numerical values that best describe this variety. The Royal Horticultural Society or any recognized color standard			
COLOR SYSTEM USED:	LOCATION OF TH	LOCATION OF THE TEST(S) TO EVALUATE THIS VARIETY:			
1. MARKET CLASS:	2 = MATURITY:	2 = MATURITY:			
CLASS 1 = Mavy (Pca) 2 = Small White 3 = Black Midnight 4 = Pinto UI-114 5 = Great Northern UI-59 6 = Small Red NW-59 7 = Pink Vive 8 = Crenberry UI-50 9 = Dark Red Kidney 10 = Light Red Kidney 11 = Yellow Eye Steuben 12 = Other (Specify	105 Da  195 Ap Note: C	populating to Harvest Maturity  Heat Units from Planting to Harvest Maturity (Optional). Specify Base Temperature Used:  ys from Planting to Harvest Maturity of Check Variety (Use Check propriate to Market Class Shown in Item 1)  alif. beans cut 2 weeks  "maturity"			
3. PLANT HABIT:		erage Height of Maluire Plant, in cm,			
1 = Is Bush-determinate, Strong and Erect Stern are 2 = Ib Bush-determinate, Weak Starn and Branches 3 = Ils Erect Growth Habit-Indeterminate, Guides ( or not developed 4 = Ilb Erect Growth Habit-Indeterminate, Guides ( with no Abitity to Climb 5 = Ills Virse-indeterminate, Short Guides with no a 6 = Illb Virse-indeterminate, Long Guides with Abitity 7 = Ive Indeterminate Climbing, Pods Distributed 1 Plant 8 = Ivb Indeterminate Climbing, Pods Concentrate Part of the Plant	Aving the Climb ty to Climb Throughout the Coding Lodging	Average Height of Check Variety, in cm. (Use Same Check as Above)  Pod Position: 1 = Low (Lower Pods Touching Soil Surface) 2 = High (Lower Pods not Touching Soil Surface) 3 = Scattered (Not Concentrated High or Low)  Adaptability to Machine Harvest: 1 = Adapted 2 = Not Adapted			





Drought

Air Pollution

Heat

12. KNOWN PHYSIOLOGICAL STRESS REACTION:

2 = Resistant

4 = Avoidance

1 = Susceptible

3 × Tolerant

Applicant: Steve TEMPLE UC Canario 707

#### **Exhibit D**

#### **Evidence in Support of Variety Distinctness Through DNA Analysis**

Amplified fragment length polymorphism (AFLP) markers were used to fingerprint yellow bean cultivars for determining the genetic relationships between yellow bean and other bean germplasm. AFLP analysis was performed as described by P. Vos et al. *Nucleic Acids Research* 23:4407 (1995), and modified by Barcaccia et al. *Plant Breed* 118:335 (1999). The primer combinations included five *Eco*RI-*Mse*I (with selective bases CAC/AAG, CAC/AGC, CCA/AGA, CCA/AGC, and CAA/AAG) and five *PstI-Mse*I combinations (AG/CAC, AG/CAT, AG/CCA, AT/CAA, and AT/CAC). The 10 AFLP primer combinations revealed 133 polymorphic amplified fragments in sampled Andean and Mesoamerican accessions.

AFLP analysis revealed that Canario 707 is found within the original Andean gene pool, whereas the cultivar "Enola" is of Peruano-descent. Canario 707 differs from Enola for 15 of 133 amplified DNA fragments, revealed by specific primers Ecac/Maag (2), Ecca/Maga (1), Ecaa/Maag (1), Pag/cac (2), Pag/Mcat (5), and Pag/Mcca (4).

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**MAY 1995** 

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# OF THE AMERICAN SOCIETY FOR HORTICULTURAL SCIENCE

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MATMAR 15 P 2: 21

J. AMER. Soc. HORT. Sci. 120(3):520-522. 1995.

# The Dark Corona Character in Seedcoats of Common Bean Cosegregates with the Pink Flower Allele $v^{\mathrm{lae}}$

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Additional index words. Phaseolus vulgaris, inheritance, genetic linkage

Abstract. Crosses were made with two common bean (Phaseolus vulgaris L.) parents that have pink flowers ( $v^{loe}/-$ ) and mineral-brown seedcoats with dark corona, viz.,  $v^{loe}$  BC, 5–593 (derived from Lamprecht V0491) and  $F_3$   $v^{loe}$  dark corona (derived from Lamprecht M0048). The third parent v BC, 5–593 had white flowers (v/v) and mineral-brown seedcoats without dark corona (derived from Lamprecht M0056). The  $F_2$  progenies of the crosses v BC, 5–593 x  $v^{loe}$  BC, 5–593 and  $F_3$   $v^{loe}$  dark corona  $\times v$  BC, 5–593 segregated for only two phenotypic classes: either pink flowers and seeds with dark corona or white flowers and seeds without dark corona. Thus, it was demonstrated that the dark corona character (Cor) is either tightly linked to  $v^{loe}$  (<4 map units) or is a pleiotropic effect of  $v^{loe}$ . Pleiotropy is more probable, but tight linkage cannot be ruled out. A linkage of 15 map units between Cor and R (currently, R is known to be tightly linked with C) reported by Lamprecht was not found by subsequent authors, and the linkage map of common bean should be revised accordingly, i.e., no linkage exists between V(Cor) and C.

In 1985 a program was initiated at the Univ. of Florida to develop genetic stocks with selected marker genes of common bean (Phaseolus vulgaris) in a common genetic background by backcrossing to a recurrent parent. The recurrent parent was a Florida dry bean breeding line, 5-593, that had been developed before 1985. One of the recessive marker characters that was backcrossed into 5-593 was the pink (laelia) flower trait controlled by the  $v^{lac}$  allele, where the dominant allele V produces bishops violet (purple) flowers and the fully recessive allele v produces pure white flowers (Lamprecht, 1935; Prakken, 1970). The genetic stock carrying the pink flower allele vlac was discovered to have a dark (black) corona around the hilum scar of the seeds, which had a mineral-brown color over the remainder of the seedcoat. The dark corona character was reported by Lamprecht (1934) to be controlled by the Cor locus. This paper describes the results of an investigation to test the hypothesis that the  $v^{lac}$  allele has an unreported pleiotropic effect on seedcoat color pattern, viz., production of a dark corona.

#### **Materials and Methods**

Florida dry bean breeding line 5–593 has determinate habit, purple flowers, and shiny, pure black seedcoats. The seedcoat genotype of 5–593 is TP [Cr] DJGBVRk (Bassett, 1994; Prakken, 1970, 1972).

Following the usual recurrent backcross procedure for developing the genetic stocks referred to above, there was strong selection in the  $F_2$  generation of each successive backcross for the phenotype of the recurrent parent. The goal was to create a genetic stock that fully recovered the appearance of 5–593 except for the selected marker trait, i.e., there was strong selection against all other marker traits. When the allele  $\nu$  was substituted into the genetic background of 5–593, giving P C D J G B  $\nu$ , it had two pleiotropic effects, viz., changing purple flowers to pure white and changing pure black seedcoats to mineral brown (Prakken, 1970). The source of the  $\nu$  allele was Lamprecht Line M0056 (now PI 527830),

Received for publication 9 Sept. 1994. Accepted for publication 14 Dec. 1994. Florida Agricultural Experiment Station Journal Series no. R-03942. The cost of publishing this paper was defrayed in part by the payment of page charges. Under postal regulations, this paper therefore must be hereby marked *advertisement* solely to indicate this fact. Professor.

which is known to carry v (Bassett et al., 1990). The resulting stock was designated v BC, 5-593, indicating two backcrosses to 5-593 with strong selection for the recurrent parent phenotype. The genetic stock vlac BC, 5-593 was created in a similar manner, using Lamprecht Line V0491 (now PI 527745) as the source of the vlac allele. When the viae allele was substituted into the genetic background of 5-593, giving P C D J G B viae, it had two wellestablished pleiotropic effects, viz., changing purple flowers to pink and changing pure black seedcoat color to mineral brown (Prakken, 1970). The resulting stock was designated vlac BC, 5-593, indicating three backcrosses to 5-593 with strong selection for the recurrent parent phenotype. However, v<sup>lae</sup> BC, 5-593 had an additional trait not reported in previous literature, viz., a dark (black) corona. The corona phenotypes of vlae BC, 5-593 (dark corona) and v BC, 5-593 (no dark corona) are illustrated in a drawing of the seedcoat patterns on the ventral side of the seeds (Fig. 1). The corona phenotype of v BC<sub>2</sub> 5-593 was slightly darker brown than the surrounding mineral-brown seedcoat and was described as having no dark corona (Fig. 1). For additional illustration and discussion of the corona character a good current reference was provided by Leakey (1988).

An additional source of pink flowers and the  $v^{\rm lac}$  allele was used in the investigation, viz., Lamprecht Line M0048 (now PI 527829). The letter M signifies multigaris in Lamprecht's terminology, i.e., a line that is derived from the interspecific cross P. vulgaris  $\times$  P. coccineus (formerly multiflorus). Line M0048 has dark seal brown seedcoat color and determinate habit. The dark seal brown color is hypothesized to be the expression of the genotype  $P[cR]JGBv^{\rm lac}$  (M.J. Bassett, unpublished data). The cross M0048  $\times$  5–593 was used to substitute the [Cr] allele of 5–593 for the [cR] allele of M0048. Thus, an  $F_3$  progeny was obtained that was true breeding for pink flowers and mineral-brown seedcoats with dark corona ( $P[Cr]JGBv^{\rm lac})$ . The line is designated  $F_3v^{\rm lac}$  dark corona.

Two crosses were made to test the hypothesis that the dark corona character is not controlled by a gene Cor that is independent of V, but is a pleiotropic effect of  $v^{\text{lae}}$ . The first was  $v \cdot BC_2 \cdot 5-593 \times v^{\text{lae}} \cdot BC_3 \cdot 5-593$ , and the second was  $F_3 \cdot v^{\text{lae}} \cdot BC_3 \cdot 5-593$ , and the second was  $F_3 \cdot v^{\text{lae}} \cdot BC_3 \cdot 5-593$ . The  $F_2 \cdot F_3 \cdot F_3$ 

color classes separate. The seeds were later examined for any indication of segregation for phenotypic classes.

I wrote letters to the curators of several common bean germplasin collections in Europe requesting seed of the two parental lines used by Lamprecht (1934) in the experiments leading to the discovery of Cor, viz., 'de la Chine' (Lamprecht Line 29) and 'Pariser Gelbe' (Lamprecht Line 23). Neither of the parental lines used by Lamprecht (1934) are extant in the U.S.Department of Agriculture plant introduction collection (at Pullman, Wash.) of Lamprecht's experimental materials, PI 527711 through PI 527878 (168 accessions). I was unsuccessful in obtaining seed of 'de la Chine', but I obtained 'Pariser Gelbe' (BGRC #25351) from the Institut für Pflanzenbau der Bundesforschungsanstalt für Landwirtschaft (FAL), Bundesallee 50, D-38116 Braunschweig, Germany. The seedcoat and corona colors of 'Pariser Gelbe' (#25351) are identical to the description of Lamprecht (1934), and the seed shape is also the same as his illustration. Seed of 'Pariser Gelbe' (#25351) was grown in the greenhouse at Gainesville and data were taken on the flower color.

#### Results and Discussion

The  $F_1$  plants from the cross  $\nu$  BC<sub>2</sub> 5–593 x  $\nu^{\rm lac}$  BC<sub>3</sub> 5–593 had pink flowers ( $\nu^{\rm lac}/\nu$ ) and mineral-brown seeds with dark corona (data not shown). The  $F_2$  segregated for only two phenotypic classes rather than the four classes that would be expected if flower color and dark corona segregated independently (Table 1). Plants with pink flower color always had dark (black) corona, and plants with white flowers always had no dark corona. Thus, there must be either tight linkage between the genes controlling the two characters or pleiotropic effects originating from the  $\nu^{\rm lac}$  allele.

There was full dominance for the dark corona trait, whereas Lamprecht (1934) reported partial dominance for dark corona. However, it should be noted that the parental materials of Lamprecht had seedcoat genotypes that were different from the parents in Table 1. For example, Lamprecht Line 23, 'Pariser Gelbe,' had genotype P C j g b v with corona, and Line 29, 'de la Chine,' had genotype P C j g b v without corona. It is probable that 'Pariser Gelbe' carried  $v^{\text{lae}}$ , but Lamprecht (1934) did not give information

on the flower color and probably omitted the superscript lae as an irrelevant distraction. This view is supported by the observation that 'Pariser Gelbe' (#25351) had pink flowers, which is characteristic of plants with  $v^{lae}$ .

The dark corona in vlae BC, 5-593 was black, whereas the corona in 'Pariser Gelbe' was a light purple color. It is my hypothesis that the action of the recessive alleles at G and B greatly reduced the quantity of anthocyannin pigment present (M.J. Bassett, unpublished data; Prakken, 1970). Therefore, the reduction in pigment produced by the heterozygote Corlcor (really vlae/v) gave a much paler purple corona (Lamprecht, 1934). The genotype cor/ cor (really v/v) had no corona (Lamprecht, 1934). The parental lines derived from 5-593 had dominant alleles at G and B and, therefore, had a much higher concentration of anthocyanin (Table 1). Thus, the heterozygote  $v^{lac}/v$  did not produce a discernible reduction in pigment, but it is probable that a reduction actually took place that is proportional to the one observed by Lamprecht (1934). It is my hypothesis that the absolute concentration of pigment must be reduced to some critical threshold level for the loss to be discernible to the naked eye. Clearly, the partial dominance of Cor (vlae) for corona expression depends on the background genotype for visual evaluation of its expression.

The F, plants from the cross F, viac dark corona x v BC, 5-593 had pink flowers and mineral-brown seed with dark corona (data not shown). In the F, progeny, only two phenotypic classes were observed, viz., plants with pink flowers and seeds with dark corona on mineral brown or plants with white flowers and seeds with no darl corona on mineral brown (Table 1). As with the previous cross, there was no independent segregation of flower color and corona color. One must conclude that either the genes are linked or the two characters are pleiotropic effects of a single locus, V. Considering the data from both crosses (Table 1), the possible linkage would be less than four map units, as estimated by adding a single (hypothetical) crossover event to the observed data and using the maximum likelihood equations of Allard (1956). One way to symbolically represent such tight linkage is to use the bracket convention, where two or more tightly linked genes are written with their gene symbols enclosed in brackets (Bassett, 1991). For the materials used in the above experiments, the

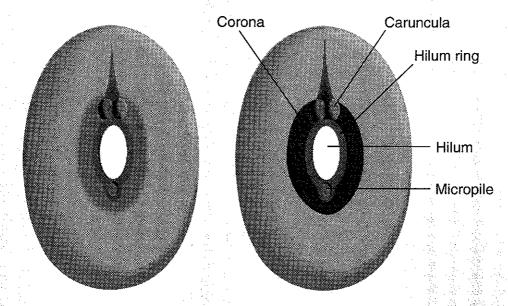


Fig. 1. The seed on the left (ν BC<sub>2</sub> 5-593) has no dark corona (light shading indicates a slightly darker brown than the surrounding mineral brown), whereas the seed on the right has dark (black) corona (ν<sup>1ω</sup> BC<sub>3</sub> 5-593).

Table 1. Segregation for flower and seed coat color in the  $F_2$  from the crosses  $\nu BC_2$ :5–593 ×  $\nu^{lae}$  BC<sub>3</sub> 5–593, and  $F_3$   $\nu^{lae}$  dark corona ×  $\nu$  BC<sub>2</sub> 5–593

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The dark corona is black in those genotypes because of dominant B. In other genetic backgrounds,  $v^{\text{lec}}$  produces various shades of purple in the corona, e.g., with recessive b or g b.

The corona is slightly darker brown than the surrounding mineral brown of the remainder of the seedcoat.

symbols would be  $[v^{lac}Cor]$  for plants with pink flowers and dark corona and [v cor] for plants with white flowers and no corona (Table 1)

The segregation for pink and white flowers in the  $F_2$  from the cross  $F_3$   $\nu^{\text{lac}}$  dark corona x  $\nu$  BC $_2$  5–593 was highly disturbed in relation to the expected 3:1 ratio for pink to white, respectively (Table 1). That result is not surprising, considering that the  $F_3$   $\nu^{\text{lac}}$  dark corona parent was derived from M0048. The initial cross M0048 x 5–593 also had a highly disturbed 3:1  $F_2$  segregation at the V locus for purple to pink flowers, respectively (data not shown). Disturbed segregation ratios in the progenies from interspecific crosses involving P. vulgaris and P. coccineus are common and have been frequently reported (as reviewed by Smartt, 1970).

Lamprecht (1961) reported a linkage of 15 map units between Cor and R and incorporated that linkage into his linkage group I. The current view is that R is tightly linked in the complex C locus (Bassett, 1991; Prakken, 1974). If Cor is tightly linked to  $v^{lae}$  as demonstrated by the data (Table 1), then the linkage of Lamprecht (1961) translates into a linkage of 15 map units between V and C. No such linkage was found in the extensive work by Nakayama (1960, 1964, 1965, 1968) and Prakken (1972) with materials segregating jointly at V and C. Linkage group I should be revised accordingly, viz., splitting linkage group I into two separate linkages: D with V and C with tri. The D locus controls hilum ring color, C controls seedcoat color, and tri controls tricotyledony (Bassett, 1991).

The data presented above are sufficient to demonstrate tight linkage between dark corona color and pink flower color, but other observations suggest that the better hypothesis is that dark corona is a pleiotropic effect of  $\nu^{\text{lae}}$ . I have observed many seedcoat

genotypes in the course of my career and have never observed either of the putative crossover phenotypes, e.g.  $(v\ Cor)$  with pure white flowers and mineral-brown seeds with dark corona due to the genotype  $TP\ CJGB\ v$ , or  $(v^{lac}\ cor)$  with pink flowers and mineral-brown seedcoats with no dark corona due to the genotye  $TP\ CJG\ B\ v^{lac}$ . Until such recombinants have been found and verified, the more probable hypothesis is pleiotropy.

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